RAW SEQUENCE LISTING

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) no errors detected.

Application Serial Number: 10/687,060 A

Source: TFWO

Date Processed by STIC: 10/25/04

ENTERED

RAW SEQUENCE LISTING   DATE: 10/25/2004
PATENT APPLICATION: **US/10/687,060A**   TIME: 15:55:22

Input Set : **A:\seqlist.txt**
Output Set: **N:\CRF4\10252004\J687060A.raw**

```
 4 <110> APPLICANT: Bruck, Claudine
 5        Godart, Stephane Andre Georges
 6        Marc-Hand, Martine
 8 <120> TITLE OF INVENTION: FUSION PROTEINS COMPRISING HIV-1 TAT
 9        AND/OR NEF PROTEINS
11 <130> FILE REFERENCE: B45110C1
13 <140> CURRENT APPLICATION NUMBER: 10/687,060A
14 <141> CURRENT FILING DATE: 2003-10-16
16 <150> PRIOR APPLICATION NUMBER: 09/509,239
17 <151> PRIOR FILING DATE: 2000-03-23
19 <150> PRIOR APPLICATION NUMBER: PCT/EP98/06040
20 <151> PRIOR FILING DATE: 1998-09-17
22 <150> PRIOR APPLICATION NUMBER: GB 9720585.0
23 <151> PRIOR FILING DATE: 1997-09-26
25 <160> NUMBER OF SEQ ID NOS: 27
27 <170> SOFTWARE: FastSEQ for Windows Version 4.0
29 <210> SEQ ID NO: 1
30 <211> LENGTH: 23
31 <212> TYPE: DNA
33 <213> ORGANISM: Artificial Sequence
35 <220> FEATURE:
36 <223> OTHER INFORMATION: PCR primer
38 <400> SEQUENCE: 1
39 atcgtccatg ggtggcaagt ggt                                 23
41 <210> SEQ ID NO: 2
42 <211> LENGTH: 23
43 <212> TYPE: DNA
44 <213> ORGANISM: Artificial Sequence
46 <220> FEATURE:
47 <223> OTHER INFORMATION: PCR primer
49 <400> SEQUENCE: 2
50 cggctactag tgcagttctt gaa                                 23
52 <210> SEQ ID NO: 3
53 <211> LENGTH: 24
54 <212> TYPE: DNA
55 <213> ORGANISM: Artificial Sequence
57 <220> FEATURE:
58 <223> OTHER INFORMATION: PCR primer
60 <400> SEQUENCE: 3
61 atcgtactag tgagccagta gatc                                24
63 <210> SEQ ID NO: 4
64 <211> LENGTH: 24
65 <212> TYPE: DNA
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```
66  <213> ORGANISM: Artificial Sequence
68  <220> FEATURE:
69  <223> OTHER INFORMATION: PCR primer
71  <400> SEQUENCE: 4
72  cggctactag tttccttcgg gcct                                    24
74  <210> SEQ ID NO: 5
75  <211> LENGTH: 23
76  <212> TYPE: DNA
77  <213> ORGANISM: Artificial Sequence
79  <220> FEATURE:
80  <223> OTHER INFORMATION: PCR primer
82  <400> SEQUENCE: 5
83  atcgtccatg gagccagtag atc                                     23
85  <210> SEQ ID NO: 6
86  <211> LENGTH: 441
87  <212> TYPE: DNA
88  <213> ORGANISM: Haemophilus influenzae
90  <400> SEQUENCE: 6
91  atggatccaa aaactttagc cctttcttta ttagcagctg gcgtactagc aggttgtagc 60
92  agccattcat caaatatggc gaatacccaa atgaaatcag acaaaatcat tattgctcac 120
93  cgtggtgcta gcggttattt accagagcat acgttagaat ctaaagcact tgcttttgca 180
94  caacaggctg attatttaga gcaagattta gcaatgacta aggatggtcg tttagtggtt 240
95  attcacgatc acttttttaga tggcttgact gatgttgcga aaaaattccc acatcgtcat 300
96  cgtaaagatg gccgttacta tgtcatcgac tttaccttaa aagaaattca aagtttagaa 360
97  atgacagaaa actttgaaac catggccacg tgtgatcaga gctcaactag tggccaccat 420
98  caccatcacc attaatctag a                                       441
100 <210> SEQ ID NO: 7
101 <211> LENGTH: 144
102 <212> TYPE: PRT
103 <213> ORGANISM: Haemophilus influenzae
105 <400> SEQUENCE: 7
106 Met Asp Pro Lys Thr Leu Ala Leu Ser Leu Leu Ala Ala Gly Val Leu
107 1               5                   10                  15
108 Ala Gly Cys Ser Ser His Ser Ser Asn Met Ala Asn Thr Gln Met Lys
109             20                  25                  30
110 Ser Asp Lys Ile Ile Ile Ala His Arg Gly Ala Ser Gly Tyr Leu Pro
111         35                  40                  45
112 Glu His Thr Leu Glu Ser Lys Ala Leu Ala Phe Ala Gln Gln Ala Asp
113     50                  55                  60
114 Tyr Leu Glu Gln Asp Leu Ala Met Thr Lys Asp Gly Arg Leu Val Val
115 65                  70                  75                  80
116 Ile His Asp His Phe Leu Asp Gly Leu Thr Asp Val Ala Lys Lys Phe
117             85                  90                  95
118 Pro His Arg His Arg Lys Asp Gly Arg Tyr Tyr Val Ile Asp Phe Thr
119             100                 105                 110
120 Leu Lys Glu Ile Gln Ser Leu Glu Met Thr Glu Asn Phe Glu Thr Met
121         115                 120                 125
122 Ala Thr Cys Asp Gln Ser Ser Thr Ser Gly His His His His His
123         130                 135                 140
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Input Set : **A:\seqlist.txt**
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```
126 <210> SEQ ID NO: 8
127 <211> LENGTH: 648
128 <212> TYPE: DNA
129 <213> ORGANISM: Human Immunodeficiency Virus
131 <400> SEQUENCE: 8
132 atgggtggca agtggtcaaa aagtagtgtg gttggatggc ctactgtaag ggaaagaatg 60
133 agacgagctg agccagcagc agatggggtg ggagcagcat ctcgagacct ggaaaaacat 120
134 ggagcaatca caagtagcaa tacagcagct accaatgctg cttgtgcctg gctagaagca 180
135 caagaggagg aggaggtggg ttttccagtc acacctcagg tacctttaag accaatgact 240
136 tacaaggcag ctgtagatct tagccacttt ttaaaagaaa aggggggact ggaagggcta 300
137 attcactccc aacgaagaca agatatcctt gatctgtgga tctaccacac acaaggctac 360
138 ttccctgatt ggcagaacta cacaccaggg ccaggggtca gatatccact gacctttgga 420
139 tggtgctaca agctagtacc agttgagcca gataaggtag aagaggccaa taaggagag 480
140 aacaccagct tgttacaccc tgtgagcctg catggaatgg atgaccctga gagagaagtg 540
141 ttagagtgga ggtttgacag ccgcctagca tttcatcacg tggcccgaga gctgcatccg 600
142 gagtacttca agaactgcac tagtggccac catcaccatc accattaa 648
144 <210> SEQ ID NO: 9
145 <211> LENGTH: 215
146 <212> TYPE: PRT
147 <213> ORGANISM: Human Immunodeficiency Virus
149 <400> SEQUENCE: 9
150 Met Gly Gly Lys Trp Ser Lys Ser Ser Val Val Gly Trp Pro Thr Val
151  1               5                  10                  15
152 Arg Glu Arg Met Arg Arg Ala Glu Pro Ala Ala Asp Gly Val Gly Ala
153             20                  25                  30
154 Ala Ser Arg Asp Leu Glu Lys His Gly Ala Ile Thr Ser Ser Asn Thr
155         35                  40                  45
156 Ala Ala Thr Asn Ala Ala Cys Ala Trp Leu Glu Ala Gln Glu Glu Glu
157     50                  55                  60
158 Glu Val Gly Phe Pro Val Thr Pro Gln Val Pro Leu Arg Pro Met Thr
159 65                  70                  75                  80
160 Tyr Lys Ala Ala Val Asp Leu Ser His Phe Leu Lys Glu Lys Gly Gly
161                 85                  90                  95
162 Leu Glu Gly Leu Ile His Ser Gln Arg Arg Gln Asp Ile Leu Asp Leu
163             100                 105                 110
164 Trp Ile Tyr His Thr Gln Gly Tyr Phe Pro Asp Trp Gln Asn Tyr Thr
165         115                 120                 125
166 Pro Gly Pro Gly Val Arg Tyr Pro Leu Thr Phe Gly Trp Cys Tyr Lys
167     130                 135                 140
168 Leu Val Pro Val Glu Pro Asp Lys Val Glu Glu Ala Asn Lys Gly Glu
169 145                 150                 155                 160
170 Asn Thr Ser Leu Leu His Pro Val Ser Leu His Gly Met Asp Asp Pro
171                 165                 170                 175
172 Glu Arg Glu Val Leu Glu Trp Arg Phe Asp Ser Arg Leu Ala Phe His
173             180                 185                 190
174 His Val Ala Arg Glu Leu His Pro Glu Tyr Phe Lys Asn Cys Thr Ser
175             195                 200                 205
176 Gly His His His His His His
177     210                 215
```

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Input Set : A:\seqlist.txt
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180 <210> SEQ ID NO: 10
181 <211> LENGTH: 288
182 <212> TYPE: DNA
183 <213> ORGANISM: Human Immunodeficiency Virus

185 <400> SEQUENCE: 10
186 atggagccag tagatcctag actagagccc tggaagcatc caggaagtca gcctaaaact 60
187 gcttgtacca attgctattg taaaaagtgt tgctttcatt gccaagtttg tttcataaca 120
188 aaagccttag gcatctccta tggcaggaag aagcggagac agcgacgaag acctcctcaa 180
189 ggcagtcaga ctcatcaagt ttctctatca aagcaacccca cctcccaatc ccgaggggac 240
190 ccgacaggcc cgaaggaaac tagtggccac catcaccatc accattaa        288

192 <210> SEQ ID NO: 11
193 <211> LENGTH: 95
194 <212> TYPE: PRT
195 <213> ORGANISM: Human Immunodeficiency Virus

197 <400> SEQUENCE: 11
198 Met Glu Pro Val Asp Pro Arg Leu Glu Pro Trp Lys His Pro Gly Ser
199  1               5                  10                  15
200 Gln Pro Lys Thr Ala Cys Thr Asn Cys Tyr Cys Lys Lys Cys Cys Phe
201             20                  25                  30
202 His Cys Gln Val Cys Phe Ile Thr Lys Ala Leu Gly Ile Ser Tyr Gly
203         35                  40                  45
204 Arg Lys Lys Arg Arg Gln Arg Arg Arg Pro Pro Gln Gly Ser Gln Thr
205     50                  55                  60
206 His Gln Val Ser Leu Ser Lys Gln Pro Thr Ser Gln Ser Arg Gly Asp
207 65                  70                  75                  80
208 Pro Thr Gly Pro Lys Glu Thr Ser Gly His His His His His
209             85                  90                  95

212 <210> SEQ ID NO: 12
213 <211> LENGTH: 909
214 <212> TYPE: DNA
215 <213> ORGANISM: Human Immunodeficiency Virus

217 <400> SEQUENCE: 12
218 atgggtggca agtggtcaaa aagtagtgtg gttggatggc ctactgtaag ggaaagaatg 60
219 agacgagctg agccagcagc agatggggtg ggagcagcat ctcgagacct ggaaaaacat 120
220 ggagcaatca caagtagcaa tacagcagct accaatgctg cttgtgcctg gctagaagca 180
221 caagaggagg aggaggtggg ttttccagtc acacctcagg tacctttaag accaatgact 240
222 tacaaggcag ctgtagatct tagccacttt ttaaaagaaa agggggggact ggaagggcta 300
223 attcactccc aacgaagaca agatatcctt gatctgtgga tctaccacac acaaggctac 360
224 ttccctgatt ggcagaacta cacaccaggg ccaggggtca gatatccact gacctttgga 420
225 tggtgctaca agctagtacc agttgagcca gataaggtag aagaggccaa taaaggagag 480
226 aacaccagct tgttacaccc tgtgagcctg catggaatgg atgaccctga gagagaagtg 540
227 ttagagtgga ggtttgacag ccgcctagca tttcatcacg tggcccgaga gctgcatccg 600
228 gagtacttca agaactgcac tagtgagcca gtagatccta gactagagcc ctggaagcat 660
229 ccaggaagtc agcctaaaac tgcttgtacc aattgctatt gtaaaaagtg ttgctttcat 720
230 tgccaagttt gtttcataac aaaagcctta ggcatctcct atggcaggaa gaagcggaga 780
231 cagcgacgaa gacctcctca aggcagtcag actcatcaag tttctctatc aaagcaaccc 840
232 acctcccaat cccgaggga cccgacaggc ccgaaggaaa ctagtggcca ccatcaccat 900
233 caccattaa                                                       909

235 <210> SEQ ID NO: 13
```

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236 <211> LENGTH: 302
237 <212> TYPE: PRT
238 <213> ORGANISM: Human Immunodeficiency Virus
240 <400> SEQUENCE: 13
241 Met Gly Gly Lys Trp Ser Lys Ser Ser Val Val Gly Trp Pro Thr Val
242  1               5                  10                  15
243 Arg Glu Arg Met Arg Arg Ala Glu Pro Ala Ala Asp Gly Val Gly Ala
244              20                  25                  30
245 Ala Ser Arg Asp Leu Glu Lys His Gly Ala Ile Thr Ser Ser Asn Thr
246          35                  40                  45
247 Ala Ala Thr Asn Ala Ala Cys Ala Trp Leu Glu Ala Gln Glu Glu Glu
248      50                  55                  60
249 Glu Val Gly Phe Pro Val Thr Pro Gln Val Pro Leu Arg Pro Met Thr
250 65                  70                  75                  80
251 Tyr Lys Ala Ala Val Asp Leu Ser His Phe Leu Lys Glu Lys Gly Gly
252                  85                  90                  95
253 Leu Glu Gly Leu Ile His Ser Gln Arg Arg Gln Asp Ile Leu Asp Leu
254              100                 105                 110
255 Trp Ile Tyr His Thr Gln Gly Tyr Phe Pro Asp Trp Gln Asn Tyr Thr
256              115                 120                 125
257 Pro Gly Pro Gly Val Arg Tyr Pro Leu Thr Phe Gly Trp Cys Tyr Lys
258          130                 135                 140
259 Leu Val Pro Val Glu Pro Asp Lys Val Glu Glu Ala Asn Lys Gly Glu
260 145                 150                 155                 160
261 Asn Thr Ser Leu Leu His Pro Val Ser Leu His Gly Met Asp Asp Pro
262              165                 170                 175
263 Glu Arg Glu Val Leu Glu Trp Arg Phe Asp Ser Arg Leu Ala Phe His
264              180                 185                 190
265 His Val Ala Arg Glu Leu His Pro Glu Tyr Phe Lys Asn Cys Thr Ser
266          195                 200                 205
267 Glu Pro Val Asp Pro Arg Leu Glu Pro Trp Lys His Pro Gly Ser Gln
268      210                 215                 220
269 Pro Lys Thr Ala Cys Thr Asn Cys Tyr Cys Lys Lys Cys Cys Phe His
270 225                 230                 235                 240
271 Cys Gln Val Cys Phe Ile Thr Lys Ala Leu Gly Ile Ser Tyr Gly Arg
272              245                 250                 255
273 Lys Lys Arg Arg Gln Arg Arg Arg Pro Pro Gln Gly Ser Gln Thr His
274          260                 265                 270
275 Gln Val Ser Leu Ser Lys Gln Pro Thr Ser Gln Ser Arg Gly Asp Pro
276          275                 280                 285
277 Thr Gly Pro Lys Glu Thr Ser Gly His His His His His His
278      290                 295                 300
281 <210> SEQ ID NO: 14
282 <211> LENGTH: 1029
283 <212> TYPE: DNA
284 <213> ORGANISM: Artificial Sequence
286 <220> FEATURE:
287 <223> OTHER INFORMATION: Fusion construct
289 <400> SEQUENCE: 14
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VERIFICATION SUMMARY   DATE: 10/25/2004
PATENT APPLICATION: **US/10/687,060A**   TIME: 15:55:23

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